WO 2005/045030 PCT/EP2004/011851

## Claims:

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1. A method for rapidly isolating nucleic acid from a nucleic acid source comprising the steps of:

- a) lysing the nucleic acid source,
  - b) filtering the lysate through a porous matrix consisting of a material based on silica or of a silica coated material to bind the nucleic acid to the porous matrix in the absence of an alcohol and in the absence of a chaotropic salt,
  - c) eluting the nucleic acid from the porous matrix of step b) by using an aqueous buffer solution.
- 2. A method according to claim 1, wherein the nucleic acid is DNA.
- 3. A method according to claim 2, wherein the DNA is genomic DNA.
- 4. A method according to claim 1 to 3, wherein the nucleic acid is of a size ranging from about 10 kbp to about 50 kbp.
- 5. A method according to claim 1, wherein the nucleic acid source is any sort of biological tissue or cell material.
  - 6. A method according to claim 5, wherein the nucleic acid source is mammalian cells, organs, biopsies, blood, serum, muscle, bone marrow, bacteria, yeast, and/or any sort of plant tissue or cells, like seeds or leaves.
  - 7. A method according to claim 1, wherein the nucleic acid source is lysed using a buffer not containing a chaotropic salt and not containing an alcohol.
- 8. A method according to claim 1, wherein a RNase and/or a protease and/or lysozyme is added to one or more of the steps of claim 1.
  - 9. A method according to claim 1, wherein the porous matrix comprises a siliceous oxide coated surface.

WO 2005/045030 PCT/EP2004/011851

10. A method according to claim 1 or 9, wherein the porous matrix is a porous silica membrane.

- 11. A method according to claim 1, 9 or 10, wherein the porous matrix comprises pores having the size ranging from 0,2  $\mu$ m to 3,2  $\mu$ m.
  - 12. A method according to claim 11, wherein the porous matrix comprises pores having the size ranging from 0,3  $\mu$ m to 2,8  $\mu$ m.
- 10 13. A method according to claim 12, wherein the porous matrix comprises pores having the size ranging from 0,5  $\mu$ m to 2,0  $\mu$ m.

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- 14. A method according to claim 1, wherein the isolated nucleic acid serves as a template in a subsequent application like AFLP, RFLP, microsatellite analysis, southern blot, PCR or quantitative real-time PCR.
- 15. A method according to claim 14, wherein the isolated nucleic acid serves as a template in a subsequent PCR or subsequent quantitative real-time PCR application.
- 16. A method according to claim 1, wherein the lysate of step a) of claim 1 is centrifuged to eliminate cell debris from the lysate prior to step b) of claim 1.
  - 17. A method according to claim 1, wherein one or more washing steps are performed subsequent to step b) of claim 1 and prior to step c) of claim 1.
  - 18. A method according to claim 17, wherein the washing step is performed using a washing buffer.
- 19. A method according to claim 1, wherein the porous matrix of step b) of claim 1 is a membrane embedded in a single column filter tube.
  - 20. A method according to claim 1, wherein the porous matrix of step b) of claim 1 is a membrane integrated in a multi-well filter plate.

WO 2005/045030 PCT/EP2004/011851

21. A method according to claims 19 and 20, wherein the membrane is assembled in one or more layers.

- 22. A method according to claim 21, wherein the pore size of one layer differs from the pore size of the other layer(s).
  - 23. A kit for performing the method according to claims 1 to 22 comprising at least:
    - a) a porous matrix consisting of a material based on silica or of a silica coated material
- b) a lysing buffer containing no alcohol and containing no chaotropic salt
  - c) an elution buffer.